5/20 #4



OIPE

RAW SEQUENCE LISTING DATE: 04/07/2003 PATENT APPLICATION: US/09/882,227 TIME: 09:47:31

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1 <110> APPLICANT: Kleanthous, Harold
         Al-Garawi, Amal
         Miller, Charles
 3
         Tomb, Jean-Francois
 1
         Ooomen, Raymond P.
 6 <120> TITLE OF INVENTION: Identification of Polynucleotides
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115	47	_		-		-				-		-		_				444
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145																		540
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180																		636
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64																		684
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67																		732
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76																		0/0
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119 120 121 123 <2: 124 <2: 125 <2: 126 <2: 127 <40 128 129 130 131	gc Al cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 Gl	t tga a agcac Q ID NGTH: PE: I GANIS QUENC t Leu u Cys	NO: 496 PRT SM: H CE: 2	ta 2 elic Ile Leu 20	obac Lys 5 Val	ctgt ter Leu Phe	ag g pylo Glu Ile	ri Lys Ile	Thre Asn	Thr 10 Lys	Phe Asp	Glu Phe	Asr.	Ala His 30	Ly: 15 Ala	s Ala	1601 1613
119 120 121 123 <2: 124 <2: 125 <2: 126 <2: 127 <40 128 129 130 131 132	gc Al cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 Gl	t tga a agcac Q ID NGTH: PE: I GANIS QUENC t Leu	NO: 496 PRT SM: H CE: 2 1 Lys	ta 2 elic Ile Leu 20	obac Lys 5 Val	ctgt ter Leu Phe	ag g pylo Glu Ile	ri Lys Ile	Thre Asn	Thr 10 Lys	Phe Asp	Glu Phe	Asr.	Ala His 30	Ly: 15 Ala	s Ala	1601 1613
119 120 121 123 <2: 124 <2: 125 <2: 126 <2: 127 <40 128 129 130 131 132 133	gc Al cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 Gl	t tga a agcac Q ID NGTH: PE: I GANIS QUENC t Lec u Cys	NO: 496 PRT EM: H CE: 2 1 Lys S Ser S Asn 35	ta 2 elic Ile Leu 20 Lys	obac Lys 5 Val Glu	ter Leu Phe	pylo Glu Ile	ri Lys Ile Glu 40	Thr Asn 25 Thr	Thr 10 Lys	Phe Asp	Glu Phe	Asr Ser Glu 45	Ala His 30 Gly	Ly: 15 Ala	s Ala Trp	1601 1613
119 120 121 123 <22 124 <22 125 <22 126 <22 127 <40 128 129 130 131 132 133 134	gc Al cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 Gl	t tga a agcac Q ID NGTH: PE: I GANIS QUENC t Leu u Cys	NO: 496 PRT EM: H CE: 2 1 Lys S Ser S Asn 35	ta 2 elic Ile Leu 20 Lys	obac Lys 5 Val Glu	ter Leu Phe	pylo Glu Ile	ri Lys Ile Glu 40	Thr Asn 25 Thr	Thr 10 Lys	Phe Asp	Glu Phe	Asr Ser Glu 45	Ala His 30 Gly	Ly: 15 Ala	s Ala Trp	1601 1613
119 120 121 123 <2: 124 <2: 125 <2: 126 <2: 127 <40 128 129 130 131 132 133	gc A1 cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 G1 Va	t tga a agcac Q ID NGTH: PE: I GANIS QUENC t Let u Cys l Lys l Phe	NO: 496 PRT EM: H CE: 2 1 Lys s Ser 35 e Leu	ta 2 elic Ile Leu 20 Lys	obac Lys 5 Val Glu Gln	ter Leu Phe Leu Glu	pylo Glu Ile Leu Asn 55	ri Lys Ile Glu 40 Lys	: Thr : Asn 25 : Thr	Thr 10 Lys Phe	Phe Asp Lys	Glu Phe Tyr Ala	Asn Ser Glu 45 Gly	Ala His 30 Gly Val	Ly: 15 Ala Gli	s Ala a Trp ı Gly s Glu	1601 1613
119 120 121 123 <2: 124 <2: 125 <2: 126 <2: 127 <40 128 129 130 131 132 133 134 135	gc A1 cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 G1 Va	t tga a agcac Q ID NGTH: PE: I GANIS QUENC t Let u Cys l Lys 50 p Asp	NO: 496 PRT EM: H CE: 2 1 Lys s Ser 35 e Leu	ta 2 elic Ile Leu 20 Lys	obac Lys 5 Val Glu Gln	ter Leu Phe Leu Glu	pylo Glu Ile Leu Asn 55	ri Lys Ile Glu 40 Lys	: Thr : Asn 25 : Thr	Thr 10 Lys Phe	Phe Asp Lys	Glu Phe Tyr Ala	Asn Ser Glu 45 Gly	Ala His 30 Gly Val	Ly: 15 Ala Gli	s Ala a Trp ı Gly s Glu	1601 1613
119 120 121 123 <2: 124 <2: 125 <2: 126 <2: 127 <40 128 129 130 131 132 133 134 135 136	gc A1 cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 G1 Va As 65	t tga a agcac Q ID NGTH: PE: I GANIS QUENC t Let u Cys l Lys 50 p Asp	CCtt NO: 496 PRT SM: H CE: 2 1 Lys S Ser 35 E Leu D Val	ta 2 elic Ile Leu 20 Lys Asp	obac Lys 5 Val Glu Gln Leu	ter Leu Phe Leu Glu Leu 70	pylo Glu Ile Leu Asn 55 Arg	ri Lys Ile Glu 40 Lys Glu	Thr 25 Thr 11e	Thr 10 Lys Phe Leu	Phe Asp Lys Tyr Cys 75	Glu Phe Tyr Ala 60 Leu	Asr Ser Glu 45 Gly Ala	Ala His 30 Gly Val	Lys 15 Als Gli Lys	s Ala Trp 1 Gly 5 Glu 7 Thi 80	1601 1613
119 120 121 123 <22 124 <22 125 <22 126 <22 127 <40 128 129 130 131 132 133 134 135 136 137	gc A1 cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 G1 Va As 65	t tga a agcac Q ID NGTH: PE: I GANIS QUENC t Let u Cys l Lys l Phe 50 p Asp	CCtt NO: 496 PRT SM: H CE: 2 1 Lys S Ser 35 E Leu D Val	ta 2 elic Ile Leu 20 Lys Asp	obac Lys 5 Val Glu Gln Leu	ter Leu Phe Leu Glu Leu 70	pylo Glu Ile Leu Asn 55 Arg	ri Lys Ile Glu 40 Lys Glu	Thr 25 Thr 11e	Thr 10 Lys Phe Leu	Phe Asp Lys Tyr Cys 75	Glu Phe Tyr Ala 60 Leu	Asr Ser Glu 45 Gly Ala	Ala His 30 Gly Val	Lys 15 Als Gli Lys	s Ala Trp 1 Gly 5 Glu 7 Thi 80	1601 1613
119 120 121 123 <22 124 <22 125 <22 126 <22 127 <40 128 129 130 131 132 133 134 135 136 137 138	gc A1 cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 G1 Va As 65 Le	t tga a agcac Q ID NGTH: PE: I GANIS QUENC t Let u Cys l Lys l Phe 50 p Asp	CCtt NO: 496 PRT SM: H CE: 2 Lys S Ser S Asn 35 Leu D Val	ta 2 elic Ile Leu 20 Lys Asp His	obac Lys 5 Val Glu Gln Leu Ala 85	ter Leu Phe Leu Glu Leu 70 Phe	pylo Glu Ile Leu Asn 55 Arg	ri Lys Ile Glu 40 Lys Glu Ser	Asn 25 Thr Ile Ser	Thr 10 Lys Phe Leu Ala Lys 90	Phe Asp Lys Tyr Cys 75 Val	Glu Phe Tyr Ala 60 Leu Gly	Asr Ser Glu 45 Gly Ala	Ala His 30 Gly Val Val	Ly: 15; Ala Gli Ly: Arc	s Ala Trp 1 Gly 5 Glu 7 Thi 80 c Cys	1601 1613
119 120 121 123 <22 124 <22 125 <22 126 <22 127 <40 128 129 130 131 132 133 134 135 136 137 138 139	gc A1 cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 G1 Va As 65 Le	t tga a a agcac Q ID NGTH: PE: I GANIS QUENC t Let Let Lys l Phe 50 p Asp	CCtt NO: 496 PRT SM: H CE: 2 Lys S Ser S Asn 35 Leu D Val	ta 2 elic Ile Leu 20 Lys Asp His	obac Lys 5 Val Glu Gln Leu Ala 85 Lys	ter Leu Phe Leu Glu Leu 70 Phe	pylo Glu Ile Leu Asn 55 Arg	ri Lys Ile Glu 40 Lys Glu Ser	Asn 25 Thr Ile Ser	Thr 10 Lys Phe Leu Ala Lys 90 Leu	Phe Asp Lys Tyr Cys 75 Val	Glu Phe Tyr Ala 60 Leu Gly	Asr Ser Glu 45 Gly Ala	Ala His 30 Gly Val Val	Ly: 15; Ala Gli Ly: Ard 95; Ala	s Ala Trp 1 Gly 5 Glu 7 Thi 80 c Cys	1601 1613
119 120 121 123 <22 124 <22 125 <22 126 <22 127 <40 128 129 130 131 132 133 134 135 136 137 138 139 140	9C A1 cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 G1 Va As 65 Le	t tga a a agcac Q ID NGTH: PE: I GANIS QUENC t Let Let Lys l Phe 50 p Asp	CCtt NO: 496 PRT EM: H CE: 2 Lys S Ser 35 E Leu D Val	ta 2 elic Ile Leu 20 Lys Asp His Leu Ser 100	obac Lys 5 Val Glu Gln Leu Ala 85 Lys	ter Leu Phe Leu 70 Phe Asp	pylo Glu Ile Leu Asn 55 Arg Lys	ri Lys Ile Glu 40 Lys Glu Ser Ala	Three Asna 25 Three Ile	Thr 10 Lys Phe Leu Ala Lys 90 Leu	Phe Asp Lys 75 Cys Val	Glu Phe Tyr Ala 60 Leu Gly	Asn Ser Glu 45 Gly Ala Val	Ala His 30 Gly Val Val Tyr Lys	Ly: 15; Ala Gli Ly: Ard 95; Ala	s Ala Trg 1 Gly 5 Glu 80 1 Cys	1601 1613
119 120 121 123 <22 124 <22 125 <22 126 <22 127 <40 128 129 130 131 132 133 134 135 136 137 138 139 140 141	9C A1 cc 10> SE 11> LE 12> TY 13> OR 00> SE Me 1 G1 Va As 65 Le	t tga a a a g C a a g C a C C C C C C C C C	CCtt NO: 496 PRT EM: H CE: 2 Lys S Ser 35 E Leu D Val	ta 2 elic Ile Leu 20 Lys Asp His Leu Ser 100 Leu	obac Lys 5 Val Glu Gln Leu Ala 85 Lys	ter Leu Phe Leu 70 Phe Asp	pylo Glu Ile Leu Asn 55 Arg Lys	ri Lys Ile Glu 40 Lys Glu Ser Ala	Thr 25 Thr Ile Ser Val Leu 105	Thr 10 Lys Phe Leu Ala Lys 90 Leu	Phe Asp Lys 75 Cys Val	Glu Phe Tyr Ala 60 Leu Gly	Asn Ser Glu 45 Gly Ala Val	Ala His 30 Gly Val Val Tyr Lys 110	Ly: 15; Ala Gli Ly: Ard 95; Ala	s Ala Trg 1 Gly 5 Glu 80 1 Cys	1601 1613

144	Asn	Lys	Lvs	Glu	Ser	Val	Leu	Lvs	Glu	Ala	Ile	Val	Ala	Leu	Glu	Leu
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146	His	Lys	Pro	Cys	Glu	Lys	Thr	Cys	Ala	Asn	Ser	Leu	Glu	Lys	Ser	Ala
147	145					150					155					160
148	Lys	Glu	Ala	Leu	Lys	Tyr	Ala	Glu	Ile	Met	Thr	Glu	Ser	Leu	Asn	Ile
149					165					170					175	
150	Val	Lys	Asp	Leu	Val	Asn	Thr	Pro	Pro	Met	Ile	Gly	Thr	Pro	Val	Tyr
151				180					185					190		
152	Met	Ala	Glu	Val	Ala	Gln	Lys	Val	Ala	Lys	Glu	Asn	His	Leu	Glu	Ile
153			195					200					205			
154		Val	His	Asp	Glu	Lys		Leu	Glu	Glu	Lys	_	Met	Asn	Ala	Phe
155		210	_				215					220	_	_	_	
156		Ala	Val	Asn	Lys		Ser	Leu	Ser	Val		Pro	Pro	Arg	Leu	
157	225	_			_	230	_	_		_	235	_	1	~ -	-	240
158	His	Leu	Val	Tyr	-	Pro	ГÀЗ	Lys	Ala	_	Lys	Lys	шe	Ala		vaı
159	~ 1	-	~ 1	-	245		.	^	01	250	.	0	т	T	255	70 T -
160	GLY	Lys	GIŻ	•	Thr	Tyr	Asp	Cys		СТА	ьеи	Ser	Leu		Pro	АТа
161	7	П	N4 - +-	260	m 1	Mat	T	7.1.	265	τ	C1	C1	C1	270	71.	W-1
162	Asp	Tyr	275	vaı	Inr	мес	ьуѕ	280	Asp	гуу	GIY	сту	285	ser	Ата	Val
163 164	Tlo	Gly		LOU	7 cn	ת ות	Lou		Lvc	Lau	Clv	V = 1		בומ	Glu	V = 1
165	116	290	пеп	цеи	ASII	ніа	295	Ата	гуэ	ьеи	Gry	300	Giu	лта	GLU	Val
166	ніс	Gly	Tla	Tla	G1 v	Δla		Glu	Aen	Mot	Tla		Pro	Δla	Δla	Tur
167	305	Сту	110	116	ОТУ	310	1111	OIU	11011	1100	315	OLY	110	7114	7114	320
168		Pro	Asp	Asp	Tle		Tle	Ser	Lvs	Glu		Lvs	Ser	Ile	Glu	
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171	_			340			-	_	345					350		
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173			355					360					365			
174	Thr	Gly	Ala	Cys	Val	Val	Gly	Leu	Gly	Glu	Phe	Thr	Ser	Ala	Ile	Met
175		370					375					380				
176	_	His	Asn	Glu	Glu		Lys	Asn	Leu	Phe		Thr	Ser	Gly	Leu	
177	385					390					395	_		_	_	400
178	Ser	Gly	Glu	Leu		Ala	Lys	Leu	Pro		Asn	Arg	His	Leu	_	Lys
179	_			_	405			_		410			_	_	415	_
180	Leu	Ile	GLu		Lys	He	Ala	Asp		Cys	Asn	ITe	Ser		Ser	Arg
181	m	C1	C1	420	т1_	m1	71 -	C1	425	Dha	T 0	7 00	C1	430	Tla	7. ~~
182	Tyr	Gly	-	Ата	ııe	Tnr	АТА		ьeu	Pne	Leu	ASII		Pne	тте	Arg
183	7 00	C1	435	T	7 00	T 110	Тхх	440	uic	т1.	7 cn	т1 ^	445	Clu	Dro	Λla
184 185	ASP	Glu 450	rne	пλ2	чэр	пλа	455	ъeu	1172	TTG	vsb	460	TTG	GTÀ	110	пта
186	ጥኒኒን	Val	Glu	Luc	Glu	Trn		Val	Asn	Ser	Pho		Δla	Ser	Glv	Δla
187	465	۷ат	ĢΙU	пуз	OLU	470	тэр	v u I	11011	JUL	475	- x y	11±U	551	OL y	480
188		Val	Ara	Ala	Cvs		Ala	Phe	Val	Glu		Leu	Leu	Lvs	Lvs	
189	O ± y	• 44	1119		485					490	Ų.I.u.		~5u	_, 5	495	
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195 196 197	<213><220><221><221><222><400>	FEAT NAMI LOCA	TURE E/KE ATIO	: Y: CI N: (DS			oylo:	ri									
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202 203 204															ggg Gly			105
205 206 207								_				_			caa Gln		-	153
208 209 210															ttg Leu			201
211 212 213		Lys	His	Gly	Trp 55	Phe	Ile	Ile	Phe	Ile 60	Gln	Arg	Tyr	Met	tat Tyr 65	Gly	Met	249
214 215 216		Arg	Thr	Ile 70	Ilė	Pro	Ile	Ser	Ile 75	Gly	Leu	Thr	Arg	Tyr 80	agc Ser	Ala	Leu	297
217 218 219		Lys	Phe 85	Āla	Ile	Ile	Asn	Leu 90	Ile	Ser	Ala	Met	Val 95	Trp	gcg Ala	Ser	Ile	345
220 221 222															cat His			393
223 224 225															tta Leu			441
226 227 228															aag Lys 145		aac Asn	489
229 230		cgc Arg	taga	agtgo	caa 1	tacaa	attci	tt ga	aaaga	atat	g aaa	attaa	aaàa	agga	agact	ttt		542
231 233 234 235	<210><211><211><212><213>	atgr SEQ LENG	ID 1 GTH: E: PI	148 RT	4		er j	pylo	ri									560
237 238 239	<400>				Gly	Leu 5	Ala	Ile	Leu	Val	Ala 10	Gly	Ile	Gly	Gly	Phe 15	Val	
240 241			Asp	Gln	Ile 20	Tyr	Phe	Tyr	Ile	Gly 25		Thr	Asn	Lys	Ala 30	-	Ile	
242 243		Gln	Lys	Lys 35		Glu	Lys	Gln	Arg 40		Lys	Leu	Ala	Leu 45	Ala	His	Leu	

Input Set : N:\vernette\US09882227.raw
Output Set: N:\CRF4\04072003\I882227.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:25; N Pos. 42,47
Seg#:27; N Pos. 920
Seq#:27; Xaa Pos. 301
Seq#:28; Xaa Pos. 301
Seq#:77; Xaa Pos. 304
Seg#:78; Xaa Pos. 304
Seq#:81; Xaa Pos. 32
Seq#:82; Xaa Pos. 32
Seq#:85; N Pos. 495
Seq#:86; Xaa Pos. 153
Seq#:103; N Pos. 71
Seq#:103; Xaa Pos. 17
Seq#:104; Xaa Pos. 17
Seq#:127; Xaa Pos. 6
Seq#:128; Xaa Pos. 6
Seq#:137; Xaa Pos. 12
Seg#:138; Xaa Pos. 12
Seq#:139; N Pos. 479,480,481,482,483
Seq#:139; Xaa Pos. 146,147,148
Seq#:140; Xaa Pos. 146,147,148
Sea#:143; N Pos. 9,12
Seq#:144; Xaa Pos. 3,4
Seg#:145; N Pos. 976,979
Seq#:145; Xaa Pos. 315,316
Seq#:146; Xaa Pos. 315,316
Seq#:153; N Pos. 409,457
Seq#:153; Xaa Pos. 128,144
Seq#:154; Xaa Pos. 128,144
Seq#:165; N Pos. 1028
Seq#:166; Xaa Pos. 326
Seq#:171; N Pos. 48
Seq#:171; Xaa Pos. 8
Seq#:172; Xaa Pos. 8
Seq#:277; Xaa Pos. 81
Seq#:278; Xaa Pos. 81
Seq#:305; N Pos. 388
Seg#:305; Xaa Pos. 118
Seq#:306; Xaa Pos. 118
Seq#:401; Xaa Pos. 217
Seg#:402; Xaa Pos. 217
Seq#:425; Xaa Pos. 182
Seq#:426; Xaa Pos. 182
Seq#:461; N Pos. 375
Seg#:461; Xaa Pos. 123
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Input Set : N:\vernette\US09882227.raw
Output Set: N:\CRF4\04072003\1882227.raw

Seq#:462; Xaa Pos. 123
Seq#:553; N Pos. 528
Seq#:553; Xaa Pos. 172
Seq#:554; Xaa Pos. 172
Seq#:585; Xaa Pos. 66
Seq#:586; Xaa Pos. 66,102
Seq#:589; Xaa Pos. 171

VERIFICATION SUMMARY DATE: 04/07/2003 PATENT APPLICATION: US/09/882,227 TIME: 09:47:32

Input Set : N:\vernette\US09882227.raw
Output Set: N:\CRF4\04072003\I882227.raw

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L:1033 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:1164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:915
M:341 Repeated in SeqNo=27
L:1214 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:288
L:4099 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:77
L:4099 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77 after pos.:963
L:4233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78 after pos.:288
L:4414 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:81
L:4414 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 after pos.:150
L:4453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82 after pos.:16
L:4649 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85 after pos.:486
L:4687 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:144
L:5530 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:103 after pos.:51
M:341 Repeated in SeqNo=103
L:5622 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:104 after pos.:16
L:6938 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:127
L:6938 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:127 after pos.:51
L:6998 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:128 after pos.:0
L:7439 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:137
L:7439 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:137 after pos.:158
L:7503 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:138 after pos.:0
L:7581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:139 after pos.:440
M:341 Repeated in SeqNo=139
L:7661 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:140 after pos.:144
L:7803 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:143 after pos.:0
L:7859 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:144 after pos.:0
L:7962 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:145 after pos.:966
M:341 Repeated in SeqNo=145
L:8014 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:146 after pos.:304
L:8269 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:153 after pos.:390
M:341 Repeated in SeqNo=153
L:8300 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:154 after pos.:112
M:341 Repeated in SeqNo=154
L:8829 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:165 after pos.:1016
L:8889 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:166 after pos.:320
L:9044 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:171 after pos.:0
M:341 Repeated in SeqNo=171
L:9106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:172 after pos.:0
L:15077 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:277
L:15077 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:277 after pos.:294
L:15149 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:278 after pos.:80
L:16693 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:305 after pos.:342
M:341 Repeated in SeqNo=305
L:16750 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:306 after pos.:112
L:22420 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:401
L:22420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:401 after pos.:675
L:22480 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:402 after pos.:208
L:23803 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:425
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/882,227 TIME: 09:47:32

DATE: 04/07/2003

Input Set : N:\vernette\US09882227.raw
Output Set: N:\CRF4\04072003\1882227.raw

L:23803 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:425 after pos.:582
L:23841 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:426 after pos.:176
L:25953 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:461 after pos.:337
M:341 Repeated in SeqNo=461
L:26065 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:462 after pos.:112
L:31225 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:537, Line#:0
L:32226 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:553 after pos.:482
M:341 Repeated in SeqNo=553
L:32272 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:554 after pos.:160
L:34202 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:585
L:34202 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:585 after pos.:243
L:34339 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ ID#:589